

SEQUENCE LISTING

<110> Martinussen, Jan
Defoor, Els Marie Celine

<120> Orotate Transporter Encoding Marker Genes

<130> 10556.204-US

<160> 22

<170> PatentIn version 3.3

 $\langle 210 \rangle$ 1

<211> 921

<212> DNA

<213> Lactococcus lactis

<220>

<221> CDS

 $\langle 222 \rangle \quad (1) \dots (921)$

<223> Orotate transporter encoding ORF

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atg tat att tac cta gct ttt gca tta gtt ggc ggt ttt tta ctt gct 48
Met Tyr Ile Tyr Leu Ala Phe Ala Leu Val Gly Gly Phe Leu Leu Ala
1 5 10 15

aat caa aat cca atc aat gcg gat tta cga aaa att gtt ggc tca cca 96
Asn Gln Asn Pro Ile Asn Ala Asp Leu Arg Lys Ile Val Gly Ser Pro
20 25 30

ttt ttg gcc tct gga att tcc aac ttt gtt ggt tcg att ttt tta gga 144
Phe Leu Ala Ser Gly Ile Ser Asn Phe Val Gly Ser Ile Phe Leu Gly
35 40 45

att atc act tta gtg acc agt caa aca ctt ttt cct agt ttt caa ttt 192
Ile Ile Thr Leu Val Thr Ser Gln Thr Leu Phe Pro Ser Phe Gln Phe
50 55 60

ggt ggc tca cac cca gta tgg ata tgg att ggt ggg gtt ctt ggt ggg 240
Val Gly Ser His Pro Val Trp Ile Trp Ile Gly Gly Val Leu Gly Gly
65 70 75 80

att ttt cta aca tct aat gtt tta ctt ttc cca aga tta gga gct gtc 288
Ile Phe Leu Thr Ser Asn Val Leu Leu Phe Pro Arg Leu Gly Ala Val
85 90 95

caa acg gtg att tta cct att ttg ggt cga ata ttg atg ggg aca ctt 336
Gln Thr Val Ile Leu Pro Ile Leu Gly Arg Ile Leu Met Gly Thr Leu
100 105 110

att gat tca ttt ggc tgg ttt cat gcc atg caa ctt ccg atg act ctg 384
Ile Asp Ser Phe Gly Trp Phe His Ala Met Gln Leu Pro Met Thr Leu
115 120 125

atg cgc ttt ttg gga gtt atc att act tta gct ggg gtt att gtc gcg Met Arg Phe Leu Gly Val Ile Ile Thr Leu Ala Gly Val Ile Val Ala 130 135 140	432
ggt ggt ctt cct aat tta aaa gaa aaa gaa gca gaa acg cac caa act Val Val Leu Pro Asn Leu Lys Glu Lys Glu Ala Glu Thr His Gln Thr 145 150 155 160	480
aac tta cta ggc tgg cga att tgg gcg gtc atc gtt ggg gca atg tcg Asn Leu Leu Gly Trp Arg Ile Trp Ala Val Ile Val Gly Ala Met Ser 165 170 175	528
gct gct caa caa gca att aat ggc aga ttg gga gtt tta ctt gaa aac Ala Ala Gln Gln Ala Ile Asn Gly Arg Leu Gly Val Leu Leu Glu Asn 180 185 190	576
act gca caa gca acc ttt gtt tcg ttc ttc att gga ttt tta gct att Thr Ala Gln Ala Thr Phe Val Ser Phe Phe Ile Gly Phe Leu Ala Ile 195 200 205	624
ttt atc gtg tct ctt ttt att gac cgc cgt ttg cca aaa att tca gaa Phe Ile Val Ser Leu Phe Ile Asp Arg Arg Leu Pro Lys Ile Ser Glu 210 215 220	672
tta aaa aaa gca aaa cct tgg aat gga att ggt gga ttt tta gga gcc Leu Lys Lys Ala Lys Pro Trp Asn Gly Ile Gly Gly Phe Leu Gly Ala 225 230 235 240	720
tca atc gtt ttt gca aca gtc gtt gct gtt ccg caa att ggt gca ggg Ser Ile Val Phe Ala Thr Val Val Ala Val Pro Gln Ile Gly Ala Gly 245 250 255	768
ctg aca att atg atg ggc ttg att gga caa att tta ggc agt atg ttg Leu Thr Ile Met Met Gly Leu Ile Gly Gln Ile Leu Gly Ser Met Leu 260 265 270	816
ggt caa caa ttt ggt tgg tgg cgc tca agt aaa tat ggc att caa att Val Gln Gln Phe Gly Trp Trp Arg Ser Ser Lys Tyr Gly Ile Gln Ile 275 280 285	864
tgg caa att gtt ggg att cta att atg ctg acc gga ata ata ttc att Trp Gln Ile Val Gly Ile Leu Ile Met Leu Thr Gly Ile Ile Phe Ile 290 295 300	912
aaa ttt tta Lys Phe Leu 305	921

<210> 2
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 <212> PRT
 <213> Lactococcus lactis

<400> 2

Met Tyr Ile Tyr Leu Ala Phe Ala Leu Val Gly Gly Phe Leu Leu Ala

1	5	10	15
Asn Gln Asn Pro Ile Asn Ala Asp Leu Arg Lys Ile Val Gly Ser Pro	20	25	30
Phe Leu Ala Ser Gly Ile Ser Asn Phe Val Gly Ser Ile Phe Leu Gly	35	40	45
Ile Ile Thr Leu Val Thr Ser Gln Thr Leu Phe Pro Ser Phe Gln Phe	50	55	60
Val Gly Ser His Pro Val Trp Ile Trp Ile Gly Gly Val Leu Gly Gly	65	70	75
Ile Phe Leu Thr Ser Asn Val Leu Leu Phe Pro Arg Leu Gly Ala Val	85	90	95
Gln Thr Val Ile Leu Pro Ile Leu Gly Arg Ile Leu Met Gly Thr Leu	100	105	110
Ile Asp Ser Phe Gly Trp Phe His Ala Met Gln Leu Pro Met Thr Leu	115	120	125
Met Arg Phe Leu Gly Val Ile Ile Thr Leu Ala Gly Val Ile Val Ala	130	135	140
Val Val Leu Pro Asn Leu Lys Glu Lys Glu Ala Glu Thr His Gln Thr	145	150	155
Asn Leu Leu Gly Trp Arg Ile Trp Ala Val Ile Val Gly Ala Met Ser	165	170	175
Ala Ala Gln Gln Ala Ile Asn Gly Arg Leu Gly Val Leu Leu Glu Asn	180	185	190
Thr Ala Gln Ala Thr Phe Val Ser Phe Phe Ile Gly Phe Leu Ala Ile	195	200	205
Phe Ile Val Ser Leu Phe Ile Asp Arg Arg Leu Pro Lys Ile Ser Glu	210	215	220
Leu Lys Lys Ala Lys Pro Trp Asn Gly Ile Gly Gly Phe Leu Gly Ala	225	230	235
			240

Ser Ile Val Phe Ala Thr Val Val Ala Val Pro Gln Ile Gly Ala Gly
245 250 255

Leu Thr Ile Met Met Gly Leu Ile Gly Gln Ile Leu Gly Ser Met Leu
260 265 270

Val Gln Gln Phe Gly Trp Trp Arg Ser Ser Lys Tyr Gly Ile Gln Ile
275 280 285

Trp Gln Ile Val Gly Ile Leu Ile Met Leu Thr Gly Ile Ile Phe Ile
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Lys Phe Leu
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 <400> 7
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 <210> 8
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 <212> DNA
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 <210> 9
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 <210> 10
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<400> 11
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